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SEQUENCE LISTING

Asn Thr Pro Val Gly Thr Asn Met Ser Tyr Glu Val Glu Ser Lys Lys

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Phe His Gln Met Pro Lys Glu Tyr Ser Gln Met Leu Gln Leu His Met

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1310

Glu Thr Ala

398

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Cys Lys Asn Leu Gln Tyr Gly Ile Glu Ser Phe Ser Asn Lys Thr Lys

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Ser Asn Pro Ala Glu Pro Val Pro Ser Ser Leu Ser Phe Leu Glu T	hr
35 40 45	

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Pro	Arg	Ala	His	Pro	Ala	Gly	Ser	Phe	His	Ala	Gly	Pro	Gly	Asp	Leu
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G1n	Lys	Trp	Ala	Gln	Ser	Gln	Asp	Gly	Phe	Glu	His	Lys	Glu	Phe	Phe
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gca cat gag ccc tcc ccg cct ggg ggc ctg gac gca cgc cag gcg ctg

Ala His Glu Pro Ser Pro Pro Gly Gly Leu Asp Ala Arg Gln Ala Leu

55

60

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Ala	Leu	Gly	Trp	Arg	Arg	Leu	Val	Pro	Ala	Val	Pro	Leu	Ser	Gln	Leu	
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							Thr									
200		10		325			1111		330	-,-	014	2,0		335	01)	
aat	222	acc	acc		cac	atc	att	aat		റമര	att	cta	acc		ccc	1231
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11311	Lys	1111	340	116	Λιg	116	116	345	Det	UIII	116	Leu	350	ASII	110	
200	oot	000		at+	as o	24+	+00		+++	000	a c.c	at a		++~	at a	1970
																1279
ser	uis	utz	rne	тте	ASP	ser	Ser	ьeu	ıyr	Lys	ASP	val	тте	Leu	val	

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			420					425					430			
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Pro	Pro	Ser	Ser	Gly	Phe	Ile	Gly	Ile	Leu	Ile	Met	Met	Ser	Met	Cys	
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Arg	Glu	Val	His	Val	Tyr	Glu	Tyr	Ile	Pro	Ser	Val	Arg	Gln	Thr	Glu	
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Leu	Cys	His	Tyr	His	Glu	Leu	Tyr	Tyr	Asp	Ala	Ala	Cys	Thr	Leu	Gly	
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Ser	Gly	Pro	Val	Gln	Pro	Lys	Gln	Lys	Leu	Leu	His	Pro	Arg	Arg	Gly
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Ala	Pro	Thr	Arg	Gly	Tyr	Glu	Lys	Asp	Val	Gly	Asn	Lys	Thr	Thr	Val
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			340					345					350		

Asp Ser Ala Leu Tyr Lys Asp Val Ile Leu Val Ala Trp Asp Pro Ala Pro Tyr Ser Ala Asn Leu Asn Leu Trp Tyr Lys Lys Pro Asp Tyr Asn Leu Phe Thr Pro Tyr Ile Gln His Arg Arg Lys Tyr Pro Thr Gln Pro Phe Tyr Ile Leu His Pro Lys Phe Ile Trp Gln Leu Trp Asp Ile Ile Gln Glu Asn Thr Arg Glu Lys Ile Gln Pro Asn Pro Pro Ser Ser Gly Phe Ile Gly Ile Leu Ile Met Met Ser Met Cys Lys Glu Val His Val Tyr Glu Tyr Ile Pro Ser Val Arg Gln Thr Glu Leu Cys His Tyr His Glu Leu Tyr Tyr Asp Ala Ala Cys Thr Leu Gly Ala Tyr His Pro Leu Leu Tyr Glu Lys Leu Leu Val Gln Arg Leu Asn Thr Gly Thr Gln Ala Asp Leu His His Lys Gly Lys Val Val Leu Pro Gly Phe Gln Thr Leu Arg Cys Pro Val Thr Ser Pro Asn Asn Thr His Ser

<210> 8

<211> 1611

<212> DNA

<213> Mouse

<400> 8

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Phe	Val	Trp	Gly	Leu	Leu	Phe	Leu	Ala	Ile	Phe	Ile	Tyr	Phe	Thr	Asr	ı	
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Ser	Asn	Pro	Ala	Ala	Pro	Met	Pro	Ser	Ser	Phe	Ser	Phe	Leu	Glu	Ser	-	
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Leu	Gln	Glu	Pro	Ser	Leu	Pro	Arg	Ser	Leu	Asp	Ala	Ser	Lys	Val	Leu	1	
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Leu	Asp	Ser	His	Pro	Glu	Asn	Pro	Phe	His	Pro	Trp	Pro	Gly	Asp	Pro	•	
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cag	aaa	tgg	gat	cag	gcc	cca	aat	ggc	ttt	gac	aat	ggg	gat	gag	ttt	33	18
Gln	Lys	Trp	Asp	Gln	Ala	Pro	Asn	Gly	Phe	Asp	Asn	Gly	Asp	Glu	Phe	•	
			100					105					110	ı			
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Phe	Thr	Ser	Gln	Val	Gly	Arg	Lys	Ser	Gln	Ser	Ala	Phe	Tyr	Pro	Glu	l	
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gaa	gat	agc	tat	ttt	ttt	gtt	gcg	gat	cag	cct	gag	ttg	tac	cac	cac	43	4
Glu	Asp	Ser	Tyr	Phe	Phe	Val	Ala	Asp	Gln	Pro	Glu	Leu	Tyr	His	His		
	130					135					140						

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tcg	aga	gcc	ttc	ctg	tac	cgg	ctc	tgg	aag	ggg	gcc	gtg	tcc	tct	aag	626
Ser	Arg	Ala	Phe	Leu	Tyr	Arg	Leu	Trp	Lys	Gly	Ala	Val	Ser	Ser	Lys	
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Pro	Leu	Val	Pro	G1y	Val	Pro	Leu	Ser	Gln	Leu	His	Pro	Arg	Gly	Leu	
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Asp	Ser	Ala	Leu	Tyr	Lys	Asp	Val	Ile	Leu	Val	Ala	Trp	Asp	Pro	Ala	
		355					360					365				
cct	tat	tct	gcc	aat	ctt	aac	ctg	tgg	tat	aag	aag	cca	gat	tac	aac	1154
Pro	Tyr	Ser	Ala	Asn	Leu	Asn	Leu	Trp	Tyr	Lys	Lys	Pro	Asp	Tyr	Asn	
	370					375					380					
ctt	ttc	act	cca	tat	atc	cag	cat	cgc	cgg	aaa	tac	ccg	act	cag	cca	1202
Leu	Phe	Thr	Pro	Tyr	Ile	G1n	His	Arg	Arg	Lys	Tyr	Pro	Thr	Gln	Pro	
385					390					395					400	
ttt	tac	att	ctt	cac	ccc	aag	ttc	ata	tgg	cag	ctt	tgg	gac	att	atc	1250
Phe	Tyr	Ile	Leu	His	Pro	Lys	Phe	Ile	Trp	Gln	Leu	Trp	Asp	Ile	Ile	
				405					410					415		
cag	gag	aat	aca	agg	gag	aag	ata	cag	ccc	aac	cca	cca	tct	tct	ggt	1298
Gln	Glu	Asn	Thr	Arg	Glu	Lys	Ile	Gln	Pro	Asn	Pro	Pro	Ser	Ser	Gly	
			420					425					430			

ttt att gga atc ctc atc atg atg tcc atg tgt aaa gag gtg cac gtg 1346 Phe Ile Gly Ile Leu Ile Met Met Ser Met Cys Lys Glu Val His Val 435 440 445 tat gag tac atc cca tct gtt cga cag aca gag ctt tgc cac tac cat 1394 Tyr Glu Tyr Ile Pro Ser Val Arg Gln Thr Glu Leu Cys His Tyr His 450 455 460 gag ctg tac tac gac gca gcc tgc acc ttg ggg gcc tac cac cca ctg 1442 Glu Leu Tyr Tyr Asp Ala Ala Cys Thr Leu Gly Ala Tyr His Pro Leu 465 470 475 480 ctc tat gaa aag cta ctg gtg cag cgc ctt aac aca ggc acc cag gca 1490 Leu Tyr Glu Lys Leu Leu Val Gln Arg Leu Asn Thr Gly Thr Gln Ala 485 490 495 gac ttg cat cac aag ggc aag gta gtc ttg cca ggc ttc cag acc ctt 1538 Asp Leu His His Lys Gly Lys Val Val Leu Pro Gly Phe Gln Thr Leu 500 505 510 cgg tgt cca gta acc agc ccc aac aat aca cat tct taa 1577 Arg Cys Pro Val Thr Ser Pro Asn Asn Thr His Ser 515 520 aatggaactc ttgggaactg atgtgcaata aggt 1611

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